

geWorkbench

geWorkbench



geWorkbench is a platform for integrated genomics, offering strong capabilities in the analysis and visualization of gene expression, sequence, and protein structure data. It offers direct access to numerous external data sources, including **caArray**, the **Cancer Gene Index** (gene-disease-compound associations), the **Cancer Gene Atlas**, **BioCarta** and the **Pathway Interaction Database**, as well as to sequence, molecular interaction, and structural databases. Written in Java for use on the desktop, geWorkbench is open-source and cross-platform, and has an extensible, component-based architecture. geWorkbench is supported by an active development effort.

For more information, visit the [geWorkbench wiki](#).